Application No:     10580922          Version No:          3.0


Input Set:


Output Set:


Started:      2008-11-24 18:42:50.065
Finished:     2008-11-24 18:42:50.689
Elapsed:      0 hr(s) 0 min(s) 0 sec(s) 624 ms
Total Warnings:    10
Total Errors:      0
No. of SeqIDs Defined:    10
Actual SeqID Count:    10

Error code     Error Description

W  402     Undefined organism found in <213>  in SEQ ID (1)

W  402     Undefined organism found in <213>  in SEQ ID (2)

W  402     Undefined organism found in <213>  in SEQ ID (3)

W  402     Undefined organism found in <213>  in SEQ ID (4)

W  402     Undefined organism found in <213>  in SEQ ID (5)

W  402     Undefined organism found in <213>  in SEQ ID (6)

W  402     Undefined organism found in <213>  in SEQ ID (7)

W  402     Undefined organism found in <213>  in SEQ ID (8)

W  402     Undefined organism found in <213>  in SEQ ID (9)

W  402     Undefined organism found in <213>  in SEQ ID (10)

<110> Barry Slobedman

     Allison Denise Abendroth

     Christina Anne Jenkins

<120> LATENT PHASE VIRAL INTERLEUKIN-10-(VII-10) AND USES THEREOF

<130> SPRUS61.001APC

<140>  10580922

<141>  2008-11-24


<150> PCT/AU2004/001675

<151> 2004-11-26


<150> AU 2003906613

<151> 2003-11-28


<160> 10


<170> PatentIn Ver. 2.1


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<212> DNA

<213> Cytomegalovirus


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tccgaggagg cgaagccggc gacgacgacg acgataaaga atacaaagcc gcagtgtcgt 180

ccagaggatt acgcgaccag attgcaagat ctccgcgtca cctttcatcg agtaaaacct 240

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tggggatgca gcgtcatgga ctggttgttg aggcggtatc tggagatcgt gttccccgca 360
ggcgaccacg tctatcccgg actcaagacg gaattgcata gtatgcgctc gacgctagaa 420
tccatctaca aagacatgcg gcaatgcgta agtgtctctg tggcggcgct gtccgcacag 480
aggtaacaac gtgttcatag cacgctgttt tactttttgtc gggctcccag cctctgttag 540
gttgcggaga taagtccgtg attagtcggc tgtctcagga ggcggaaagg aaatcggata 600
acggcacgcg gaaaggtctc agcgagttgg acacgttgtt tagccgtctc gaagagtatc 660
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Thr Lys Pro Gln Cys Arg Pro Glu Asp Tyr Ala Thr Arg Leu Gln Asp
        35                  40                  45

Leu Arg Val Thr Phe His Arg Val Lys Pro Thr Leu Gln Arg Glu Asp
    50                  55                  60

Asp Tyr Ser Val Trp Leu Asp Gly Thr Val Val Lys Gly Cys Trp Gly
 65                  70                  75                  80

Cys Ser Val Met Asp Trp Leu Leu Arg Arg Tyr Leu Glu Ile Val Phe
                85                  90                  95

Pro Ala Gly Asp His Val Tyr Pro Gly Leu Lys Thr Glu Leu His Ser
            100                 105                 110

Met Arg Ser Thr Leu Glu Ser Ile Tyr Lys Asp Met Arg Gln Cys Val
        115                 120                 125

Ser Val Ser Val Ala Ala Leu Ser Ala Gln Arg